

RAW SEQUENCE LISTING

**The Biotechnology Systems Branch of the Scientific and Technical
Information Center (STIC) no errors detected.**

Application Serial Number: 10|799,749
Source: IFWO
Date Processed by STIC: 10/14/2005

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IFWO

RAW SEQUENCE LISTING

DATE: 10/14/2005

PATENT APPLICATION: US/10/799,749

TIME: 09:40:58

Input Set : N:\Crf3\RULE60\10799749.raw

Output Set: N:\CRF4\10142005\J799749.raw

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1 <110> APPLICANT: Kreiswirth, Barry N
2   Nadich, Steven M
3 <120> TITLE OF INVENTION: System and Method for Tracking and Controlling Infections
4 <130> FILE REFERENCE: 19124.0002
5 <140> CURRENT APPLICATION NUMBER: US/10/799,749
6 <141> CURRENT FILING DATE: 2004-03-15
W--> 7 <140> CURRENT APPLICATION NUMBER: PRIOR APPLICATION NUMBER: US/10/073,256
W--> 8 <141> CURRENT FILING DATE: PRIOR FILING DATE: 2002-02-13
9 <160> NUMBER OF SEQ ID NOS: 80
10 <170> SOFTWARE: PatentIn version 3.1
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15 <213> ORGANISM: Staphylococcus aureus
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49 <212> TYPE: DNA

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50 <213> ORGANISM: Staphylococcus aureus
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57 <213> ORGANISM: Staphylococcus aureus
58 <400> SEQUENCE: 7
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64 <213> ORGANISM: Staphylococcus aureus
65 <400> SEQUENCE: 8
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69 <211> LENGTH: 24
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71 <213> ORGANISM: Staphylococcus aureus
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78 <213> ORGANISM: Staphylococcus aureus
79 <400> SEQUENCE: 10
80 gaggaagaca acaaaaaaacc tggtaaagaa gacggcaaca aacctggcaa agaagacggc 60
81 aacaagcctg gtaaagaaga caacaacaaa cctggtaaag aagacggcaa caagcctggc 120
82 aaagaagaca acaacaaacc tggcaaagaa gacggcaaca agcctggtaa agaagacaac 180
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95 <213> ORGANISM: Staphylococcus aureus
96 <400> SEQUENCE: 12
97 cgtactatcc 10
99 <210> SEQ ID NO: 13
100 <211> LENGTH: 10
101 <212> TYPE: DNA
102 <213> ORGANISM: Staphylococcus aureus
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104 attcgttata 10
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Input Set : N:\Crif3\RULE60\10799749.raw

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115 <212> TYPE: DNA	
116 <213> ORGANISM: Staphylococcus aureus	
117 <400> SEQUENCE: 15	
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121 <211> LENGTH: 10	
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129 <212> TYPE: DNA	
130 <213> ORGANISM: Staphylococcus aureus	
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135 <211> LENGTH: 12	
136 <212> TYPE: DNA	
137 <213> ORGANISM: Staphylococcus aureus	
138 <400> SEQUENCE: 18	
139 aattccccta gg	12
141 <210> SEQ ID NO: 19	
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143 <212> TYPE: DNA	
144 <213> ORGANISM: Staphylococcus aureus	
145 <400> SEQUENCE: 19	
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148 <210> SEQ ID NO: 20	
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151 <213> ORGANISM: Staphylococcus aureus	
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173 <400> SEQUENCE: 23
174      tttttcttgg caattttggt cgtattatcc gcttttttga ctgttcctga cgattcttga      60
175      tttgtctgta tctgtttagt tgcttggttt tctgctactg attcctttgt ttgactagcc      120
176      ttggcagagg gctttgaatt actttgagcg tcattaggat cttgattaga ctctaccgcg      180
177      taaatgacag aattctggcc tttgctttgg ctactttcgt ttacagtgct tggggtgcta      240
178      ctctcaactg tattgttggt tgcgctgggt gtacttgaag cactactttc gctgggtacta      300
179      cttgttttac tggttgtact tgggtgtgtg ctttcatttg tattgcttgt ttcacttgtc      360
180      gtacttgagg tactactttc gctgggtacta ctggtttcgc tggttgtgct tggcggtgtg      420
181      ctttcactcg tactactgct ctcaactgtc gtgcttggcg tgctgctttc gcttgattta      480
182      ctggtttcac ttgtcgtgct tgagggtgctg ctttcgctgg tactactgct ctcaactgtc      540
183      gtgcttggcg tgctgctttc gctgggtacta ctgctttcac ttgtcgtgct tgagggtacta      600
184      ctttcgcttg tattactggt ttcgctagtt gtacttggtg tggtgctttc atttgatttg      660
185      cttgtttcac ttgtcgtgct tgagggtgctg ctttcgcttg tattactggt tttactgggt      720
186      gtacttggtg tattgctttc atttgatttg cttgtttcac ttgttgact tgagggtgctg      780
187      ctttcgcttg tattactggt ttcactgggt gtgcttgagg tgctgctttc gcttggtgta      840
188      ctagttgtct ctgttgattt ttcaactaac agaagtaacg ccgtttttat ggtttggtgt      900
189      aattgattaa tacgcttttg tgcatctgca ggcgttttaa agccaccaag tgttggtctc      960
190      aataattctt catctgacca agcaagcagt tggtgtaact gcttagagct tccttcgcca      1020
191      gttgttgat ctattaaggc ttcttgcatg gcttgccaag agtctttggt      1070
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194 <211> LENGTH: 45
195 <212> TYPE: DNA
196 <213> ORGANISM: Enterococcus faecalis
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201 <211> LENGTH: 45
202 <212> TYPE: DNA
203 <213> ORGANISM: Enterococcus faecalis
204 <400> SEQUENCE: 25
205      gtacttgaag cactactttc gctgggtacta cttgttttac tgggtt      45
207 <210> SEQ ID NO: 26
208 <211> LENGTH: 45
209 <212> TYPE: DNA
210 <213> ORGANISM: Enterococcus faecalis
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214 <210> SEQ ID NO: 27
215 <211> LENGTH: 45
216 <212> TYPE: DNA
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Input Set : N:\Crf3\RULE60\10799749.raw

Output Set: N:\CRF4\10142005\J799749.raw

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223 <212> TYPE: DNA
224 <213> ORGANISM: Enterococcus faecalis
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226      gtgcttgagg tggtgctttc actcggtacta ctgctctcac ttgtc      45
228 <210> SEQ ID NO: 29
229 <211> LENGTH: 45
230 <212> TYPE: DNA
231 <213> ORGANISM: Enterococcus faecalis
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235 <210> SEQ ID NO: 30
236 <211> LENGTH: 45
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238 <213> ORGANISM: Enterococcus faecalis
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243 <211> LENGTH: 45
244 <212> TYPE: DNA
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250 <211> LENGTH: 45
251 <212> TYPE: DNA
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257 <211> LENGTH: 45
258 <212> TYPE: DNA
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263 <210> SEQ ID NO: 34
264 <211> LENGTH: 45
265 <212> TYPE: DNA
266 <213> ORGANISM: Enterococcus faecalis
267 <400> SEQUENCE: 34
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271 <211> LENGTH: 45
272 <212> TYPE: DNA
273 <213> ORGANISM: Enterococcus faecalis
274 <400> SEQUENCE: 35

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RAW SEQUENCE LISTING ERROR SUMMARY
PATENT APPLICATION: US/10/799,749

DATE: 10/14/2005
TIME: 09:40:59

Input Set : N:\Crf3\RULE60\10799749.raw
Output Set: N:\CRF4\10142005\J799749.raw

Invalid Line Length:

The rules require that a line not exceed 72 characters in length. This includes spaces.

Seq#:1; Line(s) 3

VERIFICATION SUMMARY

DATE: 10/14/2005

PATENT APPLICATION: US/10/799,749

TIME: 09:40:59

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Output Set: N:\CRF4\10142005\J799749.raw

L:7 M:280 W: Numeric Identifier already exists, <140> found multiple times
L:7 M:281 W: Numeric Fields not Ordered, <140> not ordered!.
L:7 M:270 C: Current Application Number differs, Replaced Current Application Number
L:8 M:280 W: Numeric Identifier already exists, <141> found multiple times
L:8 M:256 W: Invalid Numeric Header Field, Wrong Current FILING DATE:YYYY-MM-DD